# Final notebook with final function1 and model performance check function2

# model deployed in heroku

https://healthcare-prediction.herokuapp.com/

```
In [1]:
```

```
import pandas as pd
import numpy as np
import matplotlib
matplotlib.use(u'nbAgg')
import matplotlib.pyplot as plt
import warnings
warnings.filterwarnings("ignore")
import seaborn as sns
from collections import Counter
from sklearn import preprocessing
from sklearn.linear_model import LogisticRegression
from sklearn.model selection import RandomizedSearchCV
from sklearn import metrics
from sklearn.metrics import roc_curve, fl_score, confusion_matrix
from prettytable import PrettyTable
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from xgboost import XGBClassifier
import joblib
from tqdm import tqdm
```

### In [2]:

## In [22]:

```
test_inpatient = pd.read_csv('Test_Inpatientdata.csv')
test_outpatient = pd.read_csv('Test_Outpatientdata.csv')
test_bene = pd.read_csv('Test_Beneficiarydata.csv')
test_data = pd.read_csv('Test.csv')
```

# In [11]:

```
def final(X):
   """ function take raw input and then preprocess, featurize and trained """
   test inpatient = X[0]
   test_outpatient = X[1]
   test bene = X[2]
   test data = X[3]
   inpatient = pd.read csv(test inpatient)
   outpatient = pd.read csv(test outpatient)
   beneficiary = pd.read_csv(test_bene)
   test = pd.read csv(test data)
   final_data = pd.merge(inpatient, outpatient, left_on = [ col for col in outpatient.columns if c
ol in inpatient.columns], \
                   right_on = [ col for col in outpatient.columns if col in inpatient.columns], ho
= 'outer')
   final_data = pd.merge(final_data,beneficiary,how='inner',on='BeneID')
   final data = pd.merge(final data,test,how='outer',on='Provider')
   print('all file merged...')
```

```
## pre processing
    ## removing ClmProcedureCode 5, ClmProcedureCode 6 as it has all null
    final data.drop('ClmProcedureCode 5',axis=1,inplace=True)
   final data.drop('ClmProcedureCode 6',axis=1,inplace=True)
    ## remove as it has all value as same
   final data.drop('NoOfMonths PartACov',axis=1,inplace=True)
   final data.drop('NoOfMonths PartBCov',axis=1,inplace=True)
    ## replacing null with 0
   colFillna = ['ClmDiagnosisCode_1','ClmDiagnosisCode_2',
             'ClmDiagnosisCode 3', 'ClmDiagnosisCode 4',
             'ClmDiagnosisCode 5', 'ClmDiagnosisCode 6',
             'ClmDiagnosisCode_7','ClmDiagnosisCode_8',
             'ClmDiagnosisCode 9', 'ClmDiagnosisCode 10',
             'ClmProcedureCode 1', 'ClmProcedureCode 2',
             'ClmProcedureCode 3', 'ClmProcedureCode 4',
             'DiagnosisGroupCode','ClmAdmitDiagnosisCode','DeductibleAmtPaid']
    \#\# RenalDiseaseIndicator has y and 0 - replacing with 1 and 0
    final data[colFillna] = final data[colFillna].replace({np.nan:0})
    ## ChronicCond has 2 and 1 - replacing with 0 and 1
   final_data['RenalDiseaseIndicator'] = final_data['RenalDiseaseIndicator'].map(('Y':1,'0':0))
   final_data['ChronicCond_Alzheimer'] = final_data['ChronicCond_Alzheimer'].map({2:0,1:1})
   final data['ChronicCond Heartfailure'] = final data['ChronicCond Heartfailure'].map({2:0,1:1})
   final data['ChronicCond KidneyDisease'] = final data['ChronicCond KidneyDisease'].map({2:0,1:1}
   final data['ChronicCond Cancer'] = final data['ChronicCond Cancer'].map({2:0,1:1})
   final data['ChronicCond ObstrPulmonary'] = final data['ChronicCond ObstrPulmonary'].map({2:0,1:
1 } )
   final data['ChronicCond Depression'] = final data['ChronicCond Depression'].map({2:0,1:1})
   final data['ChronicCond Diabetes'] = final data['ChronicCond Diabetes'].map({2:0,1:1})
   final data['ChronicCond IschemicHeart'] = final data['ChronicCond IschemicHeart'].map({2:0,1:1}
   final data['ChronicCond Osteoporasis'] = final data['ChronicCond Osteoporasis'].map({2:0,1:1})
   final data['ChronicCond rheumatoidarthritis'] = final data['ChronicCond rheumatoidarthritis'].m
ap({2:0,1:1})
   final data['ChronicCond stroke'] = final data['ChronicCond stroke'].map({2:0,1:1})
   final data['Gender'] = final data['Gender'].map({2:0,1:1})
   print('data preprocessing done...')
    ## featurization
   ## counting num of times patient made claim
   bene count = final data['BeneID'].value counts().to dict()
   final data['BeneCount'] = final data['BeneID'].map(bene count)
    ## counting num of times provider covered claim
   pro count = final data['Provider'].value counts().to dict()
   final data['ProviderCount'] = final data['Provider'].map(pro count)
   ## counting num of times physician attended all patient
   attphy count = final data['AttendingPhysician'].value counts().to dict()
   final_data['AttendingPhysicianCount'] = final_data['AttendingPhysician'].map(attphy_count)
    ## replacing null with 0
   final data['AttendingPhysicianCount'] = final data['AttendingPhysicianCount'].fillna(0)
    ## num of days patient admitted
    ## converting non-null object type of dt column to datetime
    final data['AdmissionDt']= pd.to datetime(final data['AdmissionDt'])
    final_data['DischargeDt'] = pd.to_datetime(final_data['DischargeDt'])
    final data['ClaimStartDt']= pd.to datetime(final data['ClaimStartDt'])
    final data['ClaimEndDt']= pd.to datetime(final data['ClaimEndDt'])
    final data['numOfDaysAdmitted'] = final data['DischargeDt'] - final data['AdmissionDt']
    ## filling na with 0
   final_data['numOfDaysAdmitted'] = final_data['numOfDaysAdmitted'].fillna(0)
    ## converting days type to int
    final data['numOfDaysAdmitted'] = final data['numOfDaysAdmitted'].dt.days.astype('int64')
    ## num of days for claim took to reimbursed
```

```
final_data['numOfDaysForClaim'] = final_data['ClaimEndDt'] - final_data['ClaimStartDt']
    ## filling na with 0
   final_data['numOfDaysForClaim'] = final_data['numOfDaysForClaim'].fillna(0)
    ## converting days type to int
   final data['numOfDaysForClaim'] = final data['numOfDaysForClaim'].dt.days.astype('int64')
    ## calculating total ip,op amount reimburse
    ## adding ip op amount
   ip op total amount = final data['IPAnnualReimbursementAmt'] +
final data['OPAnnualReimbursementAmt']
   ## adding deductible amount
   ip op ded amount = final data['IPAnnualDeductibleAmt'] + final data['OPAnnualDeductibleAmt']
    ## total amount - deductible amount
   ip_op_total_amount = ip_op_total_amount - ip_op_ded_amount
   final_data['ip_op_total_amount'] = ip_op_total_amount
    ## calculating total disease patient was diagnosed before
   num of chronic = final data['RenalDiseaseIndicator'] + final data['ChronicCond Alzheimer'] + \
                    final data['ChronicCond Heartfailure'] + final data['ChronicCond KidneyDisease'
] + \
                    final data['ChronicCond Cancer'] + final data['ChronicCond ObstrPulmonary'] + \
                    final data['ChronicCond Depression'] + final data['ChronicCond Diabetes'] + \
                    final data['ChronicCond IschemicHeart'] + final data['ChronicCond Osteoporasis'
] + \
                    final data['ChronicCond rheumatoidarthritis'] + final data['ChronicCond stroke'
   final_data['num_of_chronic'] = num_of_chronic
    ## calculating num of diagnosis procedure undergonw by patient
   num of diag proc = final data[['ClmDiagnosisCode 1', 'ClmDiagnosisCode 2',
'ClmDiagnosisCode_3',
       'ClmDiagnosisCode_4', 'ClmDiagnosisCode_5', 'ClmDiagnosisCode_6',
       'ClmDiagnosisCode 7', 'ClmDiagnosisCode 8', 'ClmDiagnosisCode 9',
       'ClmDiagnosisCode_10', 'ClmProcedureCode_1', 'ClmProcedureCode_2', 'ClmProcedureCode_3', 'ClmProcedureCode_4',
       'DiagnosisGroupCode','ClmAdmitDiagnosisCode']].values
    ## counting non zero value in each row
   countnum_of_diag_proc = []
   for i in range(len(num_of_diag_proc)):
       countnum_of_diag_proc.append(np.count_nonzero(num_of_diag_proc[i]))
   final_data['num_of_diag_proc'] = countnum_of_diag_proc
    ## calculating num of physician treating patient
   num of phy = final data[['AttendingPhysician','OperatingPhysician','OtherPhysician']].fillna(0)
.values
   countnum_of_phy = []
   for i in range(len(num of phy)):
       countnum of phy.append(np.count nonzero(num of phy[i]))
   final data['num of phy'] = countnum of phy
   ## one hot encoding diagnosis code
   top10 = ['4019','25000','2724','V5869','4011','42731','V5861','2720','2449','4280']
   for col in top10:
        final_data['diagnosis_'+str(col)] = np.where(final_data['ClmDiagnosisCode_1']==col,1,0)
        final_data['diagnosis_'+str(col)] = np.where(final_data['ClmDiagnosisCode_2']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 3']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 4']==col,1,\
                                       np.where(final_data['diagnosis_'+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 5']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 6']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final_data['diagnosis_'+str(col)] = np.where(final_data['ClmDiagnosisCode_7']==col,1,\
                                       np.where(final_data['diagnosis_'+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 8']==col,1,\
                                       np.where(final_data['diagnosis_'+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 9']==col,1,\
                                      np.where(final data['diagnosis '+str(col)] == 1, 1, 0 ))
```

```
final data['diagnosis '+str(col)] = np.where(final data['ClmDiagnosisCode 10']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final data['diagnosis '+str(col)] = np.where(final data['DiagnosisGroupCode']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
        final_data['diagnosis_'+str(col)] = np.where(final_data['ClmAdmitDiagnosisCode']==col,1,\
                                       np.where(final data['diagnosis '+str(col)]==1,1,0 ))
    ## one hot encoding procedure code
   top5 = [4019.0, 9904.0, 2724.0, 8154.0, 66.0]
   for col in top5:
       final_data['procedure_'+str(col)] = np.where(final_data['ClmProcedureCode_1']==col,1,0)
       final_data['procedure_'+str(col)] = np.where(final_data['ClmProcedureCode_2']==col,1,\
                                       np.where(final data['procedure '+str(col)]==1,1,0 ))
        final data['procedure '+str(col)] = np.where(final data['ClmProcedureCode 3']==col,1,\
                                       np.where(final_data['procedure_'+str(col)]==1,1,0 ))
        final data['procedure '+str(col)] = np.where(final data['ClmProcedureCode 4']==col,1,\
                                       np.where(final_data['procedure_'+str(col)]==1,1,0 ))
    ## frequency encoding diagnosis, procedure code
   DiagnosisCode_1_count = final_data['ClmDiagnosisCode_1'].value_counts().to_dict()
   DiagnosisCode_1_count[0]=0
   DiagnosisCode_2_count = final_data['ClmDiagnosisCode_2'].value_counts().to_dict()
   DiagnosisCode 2 count[0]=0
   DiagnosisCode 3 count = final data['ClmDiagnosisCode 3'].value counts().to dict()
   DiagnosisCode 3 count[0]=0
   DiagnosisCode_4_count = final_data['ClmDiagnosisCode_4'].value_counts().to_dict()
   DiagnosisCode_4_count[0]=0
   DiagnosisCode_5_count = final_data['ClmDiagnosisCode_5'].value_counts().to_dict()
   DiagnosisCode_5_count[0]=0
   DiagnosisCode_6_count = final_data['ClmDiagnosisCode_6'].value_counts().to_dict()
   DiagnosisCode 6 count[0]=0
   DiagnosisCode_7_count = final_data['ClmDiagnosisCode_7'].value_counts().to_dict()
   DiagnosisCode_7_count[0]=0
   DiagnosisCode 8 count = final data['ClmDiagnosisCode 8'].value counts().to dict()
   DiagnosisCode_8_count[0]=0
   DiagnosisCode 9 count = final data['ClmDiagnosisCode 9'].value counts().to dict()
   DiagnosisCode 9 count[0]=0
   DiagnosisCode 10 count = final data['ClmDiagnosisCode 10'].value counts().to dict()
   DiagnosisCode 10 count[0]=0
   ClmAdmitDiagnosisCode_count = final_data['ClmAdmitDiagnosisCode'].value_counts().to_dict()
   ClmAdmitDiagnosisCode count[0]=0
   DiagnosisGroupCode count = final data['DiagnosisGroupCode'].value counts().to dict()
   DiagnosisGroupCode count[0]=0
   final data['DiagnosisCode 1 count'] =
final data['ClmDiagnosisCode 1'].map(DiagnosisCode 1 count)
   final data['DiagnosisCode 2 count'] =
final data['ClmDiagnosisCode 2'].map(DiagnosisCode 2 count)
    final_data['DiagnosisCode_3_count'] =
final data['ClmDiagnosisCode 3'].map(DiagnosisCode 3 count)
    final data['DiagnosisCode 4 count'] =
final data['ClmDiagnosisCode_4'].map(DiagnosisCode_4_count)
    final data['DiagnosisCode 5 count'] =
final_data['ClmDiagnosisCode_5'].map(DiagnosisCode_5_count)
   final_data['DiagnosisCode_6_count'] =
final data['ClmDiagnosisCode 6'].map(DiagnosisCode 6 count)
   final_data['DiagnosisCode_7_count'] =
final data['ClmDiagnosisCode 7'].map(DiagnosisCode 7 count)
    final data['DiagnosisCode 8 count'] =
final_data['ClmDiagnosisCode_8'].map(DiagnosisCode_8_count)
    final_data['DiagnosisCode_9_count'] =
final_data['ClmDiagnosisCode_9'].map(DiagnosisCode_9_count)
    final_data['DiagnosisCode_10_count'] =
final data['ClmDiagnosisCode 10'].map(DiagnosisCode 10 count)
   final data['ClmAdmitDiagnosisCode count'] = final data['ClmAdmitDiagnosisCode'].map(ClmAdmitDia
gnosisCode count)
   final data['DiagnosisGroupCode count'] = final data['DiagnosisGroupCode'].map(DiagnosisGroupCod
e count)
   print('featurization done...')
   ## dropping col after featurization
   provider = final data['Provider'].values
   final data = final data.drop(['BeneID', 'ClaimID', 'ClaimStartDt', 'ClaimEndDt', 'Provider','At
```

```
tendingPhysician', 'OperatingPhysician',
        'OtherPhysician', 'AdmissionDt', 'ClmAdmitDiagnosisCode', 'DischargeDt', 'DiagnosisGroupCode',
        'ClmDiagnosisCode_1', 'ClmDiagnosisCode_2', 'ClmDiagnosisCode_3', 'ClmDiagnosisCode_4', 'ClmDiagnosisCode_5', 'ClmDiagnosisCode_6',
        'ClmDiagnosisCode_7', 'ClmDiagnosisCode_8', 'ClmDiagnosisCode_9', 'ClmDiagnosisCode_10', 'ClmProcedureCode_1', 'ClmProcedureCode_2',
        'ClmProcedureCode 3', 'ClmProcedureCode 4', 'DOB', 'DOD'], axis=1)
     final data.to csv('final test data feature.csv')
     ## normalizing using min max scalar
    min_max_scaler = preprocessing.MinMaxScaler()
    def min max(data,column):
         """ scaling column value using min max scalar, fitting on train and transforming """
         \label{lem:min_max_scaler.fit(data[column].values.reshape(-1,1))} \\
         data scale = min max scaler.transform(data[column].values.reshape(-1,1))
         return data_scale
     ## col to normalize
    col to nor = ['InscClaimAmtReimbursed', 'DeductibleAmtPaid', 'Race',
             'State', 'County', 'IPAnnualReimbursementAmt', 'IPAnnualDeductibleAmt',
             'OPAnnualReimbursementAmt', 'OPAnnualDeductibleAmt', 'BeneCount',
             'ProviderCount', 'AttendingPhysicianCount', 'numOfDaysAdmitted',
'numOfDaysForClaim', 'ip_op_total_amount', 'num_of_chronic',
'num_of_diag_proc', 'num_of_phy', 'DiagnosisCode_1_count', 'DiagnosisCode_2_count',
             'DiagnosisCode_3_count', 'DiagnosisCode_4_count',
             'DiagnosisCode_5_count', 'DiagnosisCode_6_count',
             'DiagnosisCode_7_count', 'DiagnosisCode_8_count',
             'DiagnosisCode_9_count', 'DiagnosisCode_10_count',
             'ClmAdmitDiagnosisCode count', 'DiagnosisGroupCode count']
    for i in col to nor:
         data scale = min max(final data,i)
         final data[i] = data scale
    print('normalizing done...')
     ## loading model
    def predict using best threshold(probability, threshold):
         """ convert proba value using best threshold """
         predictions = []
         for i in probability:
              if i>=threshold:
                  predictions.append(1)
                 predictions.append(0)
         return predictions
    model = joblib.load('xgb_tuned_on_0.pkl')
    print('loading model done...')
    y_pred = model.predict_proba(final_data)[:,1]
    pred on best threshold = predict using best threshold(y pred, threshold = 0.2092)
    prediction = pd.DataFrame()
    prediction['providerID'] = provider
    prediction['predictions'] = pred_on_best_threshold
    print('prediction done...')
    return prediction
In [12]:
prediction on test= final(X)
print(prediction on test)
all file merged...
data preprocessing done...
featurization done...
normalizing done...
loading model done...
prediction done...
        providerID predictions
          PRV57070
```

1

PRV57070

1

```
2
      PRV57070
                         0
3
      PRV57070
                         1
4
      PRV57070
                        1
      PRV57070
5
                         1
       PRV57070
                         1
6
      PRV57070
7
                         1
      PRV57070
8
                        1
      PRV57070
10
      PRV57070
                        1
      PRV57070
11
                        1
12
       PRV56558
                         1
      PRV56558
1.3
                         1
14
      PRV56558
                         1
15
      PRV56558
                        1
      PRV56558
16
                         1
17
       PRV56558
                         1
      PRV56558
18
                         1
      PRV56558
19
                         1
20
      PRV56558
      PRV56558
21
                        1
22
       PRV56558
                        1
23
       PRV56558
                         1
      PRV56558
24
                         1
25
      PRV56558
                        1
26
      PRV56558
      PRV56558
27
                        1
28
       PRV56558
                         1
      PRV56558
29
                         1
                       . . .
135362 PRV53737
                       0
135363 PRV53737
      PRV51659
                        0
135364
135365
       PRV51659
                        0
135366 PRV51659
                        0
135367 PRV51659
                        0
135368 PRV53557
                        0
                        0
135369 PRV54791
135370
       PRV54791
                        0
      PRV52620
135371
                        0
135372 PRV52620
                        0
135373 PRV56192
                        0
135374 PRV56192
      PRV51728
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135375
135376
       PRV51728
                        0
135377
      PRV51728
                        0
135378 PRV56619
135379 PRV56082
                        0
                        0
135380 PRV54169
135381
       PRV56325
      PRV57628
135382
                        0
135383 PRV52776
                        0
135384 PRV54613
135385 PRV56372
                        0
135386
       PRV56372
                        0
135387
       PRV54617
                        0
135388 PRV53736
                        0
135389 PRV53736
135390 PRV53089
                        0
135391 PRV55336
```

[135392 rows x 2 columns]

## In [14]:

```
## file with null filled with 0

final_data_test = pd.read_csv('final_data_feature.csv')
final_data_test.drop('Unnamed: 0',axis=1,inplace=True)
```

# In [15]:

```
from sklearn.model_selection import train_test_split
train, test = train_test_split(final_data_test, test_size=0.20,random_state=0)
```

```
In [17]:
Y = test['PotentialFraud']
X = test.drop('PotentialFraud', axis=1)
In [19]:
X.shape
Out[19]:
(111643, 58)
In [23]:
def performance(X,Y):
    """ take input test and predict on trained model and check model performance"""
    ## normalizing using min max scalar
    min max scaler = preprocessing.MinMaxScaler()
    def min max(data,column):
         """ scaling column value using min max scalar, fitting on train and transforming """
        min max scaler.fit(data[column].values.reshape(-1,1))
        data_scale = min_max_scaler.transform(data[column].values.reshape(-1,1))
        return data scale
    ## col to normalize
    col to nor = ['InscClaimAmtReimbursed', 'DeductibleAmtPaid', 'Race',
            'State', 'County', 'IPAnnualReimbursementAmt', 'IPAnnualDeductibleAmt',
            'OPAnnualReimbursementAmt', 'OPAnnualDeductibleAmt', 'BeneCount',
            'ProviderCount', 'AttendingPhysicianCount', 'numOfDaysAdmitted', 'numOfDaysForClaim', 'ip_op_total_amount', 'num_of_chronic',
            'num of diag proc', 'num of phy', 'DiagnosisCode 1 count', 'DiagnosisCode 2 count',
            'DiagnosisCode_3_count', 'DiagnosisCode_4_count', 'DiagnosisCode_5_count', 'DiagnosisCode_6_count',
            'DiagnosisCode 7 count', 'DiagnosisCode 8 count', 'DiagnosisCode 9 count', 'DiagnosisCode 10 count',
            'ClmAdmitDiagnosisCode_count', 'DiagnosisGroupCode_count']
    for i in col to nor:
         data_scale = min_max(X,i)
        X[i] = data_scale
     ## loading model
    def predict using best threshold(probability, threshold):
         """ convert proba value using best threshold """
        predictions = []
         for i in probability:
             if i>=threshold:
                 predictions.append(1)
             else:
                predictions.append(0)
         return predictions
    def plot confusion matrix(model, X, Y, threshold):
         """ ploting confusion matrix using best threshold """
         ## fpr,tpr,threshold
        y pred = model.predict proba(X)[:,1]
        fpr,tpr,thresholds = roc curve(Y, y pred)
         ## confusion matrix
        test conmat = confusion matrix(Y, predict using best threshold(y pred, threshold))
        sns.heatmap(test conmat,xticklabels=['Predicted No','Predicted Yes'],yticklabels=['Actual N
o', 'Actual Yes'], annot=True, fmt='d')
        plt.title('confusion matrix on test')
        plt.show()
    def f1 (model, X, Y, threshold):
         """ calculate f1, macro f1 score """
         y pred = model.predict_proba(X)[:,1]
         fpr,tpr,thresholds = roc curve(Y, y pred)
         f1 test = f1 score(Y, predict using best threshold(y pred, threshold))
```

```
macro_fl_test = fl_score(Y, predict_using_best_threshold(y_pred, threshold), average='macro'

return fl_test, macro_fl_test

model = joblib.load('xgb_tuned_on_0.pkl')
    print('loading model done...')

y_pred = model.predict_proba(X)[:,1]
    pred_on_best_threshold = predict_using_best_threshold(y_pred, 0.2092)

plot_confusion_matrix(model,X,Y,0.2092)
    fl,macro_fl = fl(model,X,Y,0.2092)
    print('fl score :',fl)
    print('macro_fl score :',macro_fl)

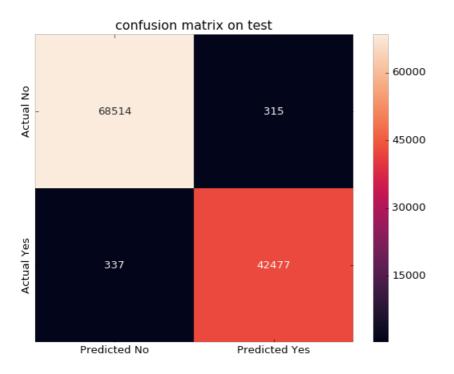
prediction = pd.DataFrame()
    prediction['actual_Y'] = Y
    prediction['predictions'] = pred_on_best_threshold
    print('prediction done...')

return prediction
```

#### In [24]:

```
model_performance = performance(X,Y)
print(model_performance)
```

loading model done...



```
fl score: 0.992383711422
macro f1 score : 0.993824046298
prediction done...
       actual_Y predictions
368022
              0
                           0
347889
              0
                            0
6731
              1
                            1
371099
              0
                            0
487726
              0
                            0
270660
              1
                            1
453174
              0
                            0
420316
              0
                            0
334583
              0
                            0
```

319996 120196 150101 204402 6665 210765 76439 128651 119971 268372 433588 287115 99422 205597 212789 415365 398477 432463 336770 473996 433572	1 0 0 0 1 1 1 0 1 1 0 1 0 1 0 0 1 1 0 0 1	1 1 0 0 1 1 1 0 1 1 0 0 1 1 0 0 1 1 0 0 0 1 1 0
125085 516734 495158 450271 294811 436077 32293 367910 415508 393524 279290 443780 101359 501260 300732 297609 483147 151583 310379 21337 38649 422069 140751 390808 299621 538252 216964 36254 386387 119136	1 0 0 0 0 0 0 1 0 0 0 1 1 0 0 0 1 1 0 0 0 1 1 0 0 1 1 0 0 0 1 1 0	1 0 0 0 0 0 0 1 1 0 0 0 1 1 1 0 0 0 1 1 1 0 0 0 1 1 1 0

[111643 rows x 2 columns]

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